

GenCore version 6.3
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OM nucleic - nucleic search, using sw model

Run on: June 26, 2009, 03:01:55 ; Search time 109 Seconds
(without alignments)
20559.273 Million cell
updates/sec

Title: US-09-300-482-225
Perfect score: 316
Sequence: 1
gataagggtgcgcacactgag.....gaatggagggagtacttcga 316

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 9875436 seqs, 3558593875 residues

Total number of hits satisfying chosen parameters: 19750872

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA:*

1: /ABSS/Data/CRF/ptodata/1/ina/1_COMB.seq:*

2: /ABSS/Data/CRF/ptodata/1/ina/5_COMB.seq:*

3: /ABSS/Data/CRF/ptodata/1/ina/6A_COMB.seq:*

4: /ABSS/Data/CRF/ptodata/1/ina/6B_COMB.seq:*

5: /ABSS/Data/CRF/ptodata/1/ina/7A_COMB.seq:*

6: /ABSS/Data/CRF/ptodata/1/ina/7B_COMB.seq:*

7: /ABSS/Data/CRF/ptodata/1/ina/7C_COMB.seq:*

8: /ABSS/Data/CRF/ptodata/1/ina/7D_COMB.seq:*

9: /ABSS/Data/CRF/ptodata/1/ina/HA_COMB.seq:*

10: /ABSS/Data/CRF/ptodata/1/ina/HB_COMB.seq:*

11: /ABSS/Data/CRF/ptodata/1/ina/PCTUS_COMB.seq:*

12: /ABSS/Data/CRF/ptodata/1/ina/PP_COMB.seq:*

13: /ABSS/Data/CRF/ptodata/1/ina/RE_COMB.seq:*

14: /ABSS/Data/CRF/ptodata/1/ina/backfiles1.seq:*